



#4

## SEQUENCE LISTING

<11> Ditzel, H.  
<12> Baden, D.  
<13> Schaller, M.

<120> Autoantibodies to glucose-6-phosphate isomerase and their participation in autoimmune disease

<130> 1361.005US1

<140> US 09/828,708

<141> 2001-04-06

<160> 123

<170> FastSEQ for Windows Version 4.0

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<213> Homo sapiens

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Ala Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr  
35 40 45  
Trp Ala Ser Thr Arg Glu Ser Gly Val Pro Asp Arg Phe Ser Gly Ser  
50 55 60  
Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Ala Glu  
65 70 75 80  
Asp Val Ala Val Tyr Tyr Cys Gln Gln Tyr Tyr Asp Ser Tyr Thr Phe  
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Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Thr Val Ala  
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<211> 104

<212> PRT

<213> Homo sapiens

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Pro Gly Lys Ala Pro Lys Leu Leu Ile Tyr Ala Ala Ser Thr Leu Gln  
35 40 45  
Ser Gly Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Glu Phe  
50 55 60  
Thr Leu Thr Ile Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr  
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Val Gly Ile Arg Arg Thr Val Ala  
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Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr Gly Ala Ser Ser Arg  
35 40 45  
Ala Thr Gly Ile Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp  
50 55 60  
Phe Thr Leu Thr Ile Ser Arg Leu Glu Pro Glu Asp Phe Ala Val Tyr  
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Tyr Cys Gln Gln Tyr Gly Ser Ser Pro Arg Thr Phe Gly Gln Gly Thr  
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Lys Val Glu Ile Lys Arg Thr Val Ala  
100 105

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Lys Ser Ser Gln Ser Val Phe Tyr Thr Ser Asn Asn Lys Asn Tyr Leu  
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Ala Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr  
35 40 45  
Trp Ala Ser Thr Arg Glu Ser Gly Val Pro Asp Arg Phe Ser Gly Ser  
50 55 60  
Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Ala Glu  
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35 40 45  
Ala Thr Gly Ile Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp  
50 55 60  
Phe Ser Phe Thr Ile Ser Ser Leu Gln Pro Glu Asp Thr Gly Thr Tyr

65                   70                   75                   80  
Tyr Cys Gln Gln Tyr Asp Asn Val Pro Asp Thr Phe Gly Gln Gly Thr  
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Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr Gly Ala Ser Ser Arg  
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Ala Thr Gly Ile Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp  
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Phe Thr Leu Thr Ile Ser Arg Leu Glu Pro Glu Asp Phe Ala Val Tyr  
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    20                25                30  
Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr Gly Ala Ser Ser Arg  
    35                40                45  
Ala Thr Gly Ile Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp  
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Phe Thr Leu Thr Ile Ser Arg Leu Glu Pro Glu Asp Phe Ala Val Tyr  
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Tyr Cys Gln Gln Tyr Gly Ser Ser Pro Arg Thr Phe Gly Gln Gly Thr  
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Lys Val Glu Ile Lys Arg Thr Val Ala  
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    20                25                30

Ala Pro Gly Lys Gly Leu Glu Trp Val Ala Leu Leu Ser Ser Asp Gly  
35 40 45  
Ser Asn Lys Phe Tyr Ile Glu Ser Val Lys Gly Arg Phe Thr Ile Ser  
50 55 60  
Lys Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg  
65 70 75 80  
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85 90 95  
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<210> 9  
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<212> PRT  
<213> Homo sapiens

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Ala Pro Gly Lys Gly Leu Glu Trp Val Ala Leu Leu Thr Met Asp Arg  
35 40 45  
Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln Leu  
50 55 60  
Ser Ser Leu Arg Pro Glu Asp Thr Ala Val Tyr Tyr Cys Thr Asn Ser  
65 70 75 80  
Glu Val Gly Ala Thr Ala Phe Asp Tyr Trp Gly Gln Gly Thr Leu Val  
85 90 95  
Thr Val Ser Ser  
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<212> PRT  
<213> Homo sapiens

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Ala Pro Gly Lys Gly Leu Glu Trp Val Ala Val Ile Ser Tyr Asp Gly  
35 40 45  
Asn Lys Lys Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser  
50 55 60  
Lys Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg  
65 70 75 80  
Val Glu Asp Thr Ala Val Tyr Tyr Cys Ala Ile Ser Ile Val Gly Thr  
85 90 95  
Thr Ala Phe Asn Tyr  
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<212> PRT  
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<400> 11

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Ala Pro Gly Gln Gly Leu Gln Trp Met Gly Arg Ile Asn Pro Thr Gly  
35 40 45  
Gly Gly Val Ser Leu Ala Gln Ser Phe Gln Asp Arg Val Ser Leu Thr  
50 55 60  
Arg Asp Arg Ser Ser Asn Thr Val Phe Leu Glu Leu Ser Gly Leu Thr  
65 70 75 80  
Glu Glu Asp Thr Ala Leu Tyr Phe Cys Ala Arg Pro Arg Phe Asn Met  
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Ile Arg Glu Pro Leu Asp Leu Trp Gly Gln Gly Thr Val Val Thr Val  
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Ala Pro Gly Lys Gly Leu Glu Trp Val Ser Arg Ile Ser Gly Asn Ser  
35 40 45  
Gly Ser Thr Phe Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser  
50 55 60  
Arg Asp Asn Ser Lys Asn Thr Ala Phe Leu Arg Met Asn Ser Gln Arg  
65 70 75 80  
Ala Glu Asp Thr Ala Val Tyr Tyr Cys Ala Lys Asp Leu Ser Ser Gly  
85 90 95  
Ala Tyr Tyr Tyr Tyr Gly Met Asp Val Trp Gly Gln Gly Thr Thr Val  
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Thr Val Ser Ser  
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Gly Pro Gly Leu Val Arg Pro Ser Gln Thr Leu Ser Leu Thr Cys Pro  
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20 25 30  
Arg Gln Pro Val Gly Lys Gly Leu Glu Trp Ile Gly Arg Ile Tyr Gly  
35 40 45  
Arg Gly Thr Thr Asn Tyr Asn Arg Val Phe Gly Ser Arg Val Ser Met  
50 55 60  
Ser Val Asp Met Ser Arg Ser Gln Phe Phe Leu Glu Leu Arg Asp Val  
65 70 75 80  
Thr Ala Ala Asp Thr Ala Val Tyr Tyr Cys Ala Arg Asp Lys Gly Ser

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20                 25                 30  
Ala Pro Gly Gln Gly Leu Glu Trp Met Gly Gly Ile Ile Pro Pro Phe  
35                 40                 45  
Gly Pro Val Asn Tyr Ala Gln Lys Phe Gln Gly Arg Val Thr Ile Thr  
50                 55                 60  
Ala Asp Asp Ser Thr Asn Thr Ala Tyr Met Gly Leu Ser Ser Leu Arg  
65                 70                 75                 80  
Ser Gly Asp Thr Ala Val Tyr Tyr Cys Ala Arg Val Ala Tyr Asp Gly  
85                 90                 95  
Ser Gly Tyr Tyr Asn Asn Ile Pro Lys Ile Tyr Tyr Tyr Ser Tyr Met  
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Gly

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Asp

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Arg Ile Ser Gly Asn Ser Gly Ser Thr Phe Tyr Ala Asp Ser Val Lys  
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Gly

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Tyr Tyr Ser Tyr Met Asp Val  
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Arg Ala Ser Gln Gly Ile Ser Ser Tyr Leu Ala  
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Gln Gln Tyr Gly Ser Ser Pro Arg Thr  
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Ala Ser Gly Phe Thr Phe Ser  
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Ala Ser Gly Asn Thr Phe Thr  
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Gly Gly Gly Leu Val Gln Pro Gly Gly Ser Leu Arg Leu Ser Cys Ala  
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Thr Ser Gly Phe Ile Phe Asn  
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<400> 62  
Gly Pro Gly Leu Val Arg Pro Ser Gln Thr Leu Ser Leu Thr Cys Pro  
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Val Ser Pro Gly Ser Ile Lys  
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Trp Val Arg Gln Pro Val Gly Lys Gly Leu Glu Trp Ile Gly  
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Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met Gly  
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<400> 69

Arg Phe Thr Ile Ser Lys Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln  
1 5 10 15  
Met Asn Ser Leu Arg Ile Asp Asp Thr Ala Val Tyr Tyr Cys Ala Ile  
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<400> 70

Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln  
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Leu Ser Ser Leu Arg Pro Glu Asp Thr Ala Val Tyr Tyr Cys Thr Asn  
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Arg Phe Thr Ile Ser Lys Asp Asn Ser Lys Asn Thr Leu Tyr Leu Gln  
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Met Asn Ser Leu Arg Val Glu Asp Thr Ala Val Tyr Tyr Cys Ala Ile  
20 25 30

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<400> 72  
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20 25 30

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<400> 73  
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20 25 30

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20 25 30

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Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser  
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Trp Gly Gln Gly Ile Val Val Asn Val Phe Ser  
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<211> 16

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<213> Homo sapiens

<400> 81

Pro Asp Ser Leu Ala Val Ser Leu Gly Glu Arg Ala Thr Ile Asn Cys  
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<211> 16

<212> PRT

<213> Homo sapiens

<400> 82

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<211> 16

<212> PRT

<213> Homo sapiens

<400> 83

Pro Gly Thr Leu Ser Leu Ser Pro Gly Glu Arg Ala Thr Leu Ser Cys  
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<210> 84

<211> 16

<212> PRT

<213> Homo sapiens

<400> 84

Pro Asp Ser Leu Ala Val Ser Leu Gly Glu Arg Ala Thr Ile Asn Cys  
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<213> Homo sapiens

<400> 86

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<211> 16

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<213> Homo sapiens

<400> 87

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<213> Homo sapiens

<400> 88

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<210> 89

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<212> PRT

<213> Homo sapiens

<400> 89

Trp Tyr Gln Leu Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile Tyr  
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<211> 15

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<213> Homo sapiens

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<400> 91  
Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr  
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<210> 92  
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<400> 92  
Trp Tyr Gln Gln Arg Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr  
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<210> 93  
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<400> 93  
Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Arg Leu Leu Ile Tyr  
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<210> 94  
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<400> 94  
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<210> 95  
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<400> 95  
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<210> 96  
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<400> 96  
Gly Val Pro Ser Arg Phe Ser Gly Ser Gly Thr Glu Phe Thr

1 5 10 15  
Leu Thr Ile Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys  
20 25 30

<210> 97  
<211> 32  
<212> PRT  
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<400> 97  
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Leu Thr Ile Ser Arg Leu Glu Pro Glu Asp Phe Ala Val Tyr Tyr Cys  
20 25 30

<210> 98  
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<212> PRT  
<213> Homo sapiens

<400> 98  
Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr  
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Leu Thr Ile Ser Ser Leu Gln Ala Glu Asp Val Ala Val Tyr Tyr Cys  
20 25 30

<210> 99  
<211> 32  
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<400> 99  
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Phe Thr Ile Ser Ser Leu Gln Pro Glu Asp Thr Gly Thr Tyr Tyr Cys  
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20 25 30

<210> 101  
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<400> 101  
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<400> 102  
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<210> 103  
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<400> 103  
Phe Gly Gly Gly Ala Lys Val Gly Ile Arg Arg Thr Val Ala  
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<400> 104  
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1 5 10

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<212> PRT  
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Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Thr Val Ala  
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 cactttgtc gtctgatgga agtaataaat tctatataga atccgtgaag ggcgcattca 180  
 ccatctccaa ggacaattct aagaacacac tgtatctgca aatgaacacgc ctgagaattg 240  
 aegacacggc tgtctattac tgtgcgattt ccctgggtggg aactaccgct tttaactact 300  
 ggggccaggg aaccctggtc accgtctcct ca 332

<210> 110  
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 <212> DNA  
 <213> Homo sapiens

<400> 110  
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 agtagtcata ccatgcactg ggtccggccag gctccaggca aggggctgga gtgggtggca 120  
 ctatattct atgatggaag taataaaatc tatcagact ccgtgaaggg ccgattcacc 180  
 atctccagag acaattccaa gaacacgctg tatctgcaat tgagcaggct aagacctgag 240  
 gacacggctg tctattattg tacgaattcc gaggtgggag ctaccgctt tgactactgg 300  
 ggccagggaa ccctggtcac cgtctcctca g 331

<210> 111  
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 <212> DNA  
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<400> 111  
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 accttcagtt cctatacttt ccactgggtc cgccaggctc caggcaaggg gctggagtgg 120  
 gtggcagtt tatcatatga tggaaacaag aaatactacg cagactccgt gaagggccga 180  
 ttcaccatct ccagagacaa ttccaagaac acttataatc tgcaaataaa cagcctgaga 240  
 gttgaggaca cggctgttta ttactgtgcg atttccatag tgggaactac cgcttttaac 300  
 tactggggcc agggAACCTT ggtcaccggtc tcctc 335

<210> 112  
 <211> 327  
 <212> DNA  
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 agtttttt acacttccaa caataagaac tacttagttt ggtaccagca gaaaccaggc 120  
 cagcctccta agttgctcat ttactggca tccacccggg aatccgggtt ccctgaccga 180  
 ttcagtggca gcgggtctgg gacagatttc acttcacca tcagcaggct gcaggctgaa 240  
 gatgtggcag tttattactg tcagcaatat tatgattcgt acactttgg ccaggggacc 300  
 aagctggaga tcaaacgaac tgtggct 327

<210> 113  
 <211> 312

<212> DNA  
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<400> 113  
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 ggcatttagca gttatttagc ctggtatcatcg ctaaaaccgg gaaaagcccc taagctcctg 120  
 atctatgtcg catccacttt gcaaagtggg gtcccatcaa gggtcagcgg cagtgatct 180  
 gggacagaat tcactctcac aataaggcgc ctgcagcctg aagatttgc aacttattac 240  
 tgtcaacagc ttaatagttt ccctctcaact ttcggcggag gggccaaggt ggggatcaga 300  
 cgaactgtgg ct 312

<210> 114  
 <211> 315  
 <212> DNA  
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 ctcatctatg gtgcataccag cagggccact ggcataccag acaggttcag tggcagtgg 180  
 tctgggacag acttcactct caccatcagc agactggagc ctgaagattt tgcaagtat 240  
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<210> 115  
 <211> 327  
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 cagccctcta agttgctcat ttactggca tccacccggg aatccggggtt ccctgaccga 180  
 ttcagtggca gcgggtctgg gacagatttc actctcacca tcagcagcct gcaggctgaa 240  
 gatgtggcag tttattactg tcagcaatat tatgattcgt acactttgg ccaggggacc 300  
 aagctggaga tcaaacgaac tggct 327

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 <212> DNA  
 <213> Homo sapiens

<400> 116  
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 ctcatctatg gtgcataccag cagggccacc ggcataccag acagatttcag tggaaagtgg 180  
 tctgggacag atttcagttt caccatcagc agtctgcagc ctgaagatac tgggacat 240  
 tactgtcaac aatatgataa tggcttcgtc acctttggcc agggggacca gctggagatc 300  
 aacgaactg tggct 315

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 <213> Homo sapiens

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ctcatctatg gtgcacccag tagggccact ggcacccag acaggttcag tggcagtgg	180
tctggacag acttcactct caccatcagc agactggagc ctgaagattt tgcagtgat	240
tactgtcagc agtatggtag ctcacccctc ttccggccaag ggacacgact ggagattaaa	300
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ctcatctatg gtgcacccag cagggccact ggcacccag acaggttcag tggcagtgg	180
tctggacag acttcactct caccatcagc agactggagc ctgaagactt tgcagttat	240
tactgtcagc agtatggaaag ctcacccctc acgttcggcc aagggaccaa ggtggaaatc	300
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<211> 342	
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atgggaagaa tcaaccccgac tggcggccggc gtttgtctcg cacagatgtt ccaggacaga	180
gtcagcctga ccagggacag gtcgtccaat acagtcttct tggaaactgag cggcctcact	240
gaggaggaca cggccttata tttctgtgcg aggccccat ttaacatgtat ccgggaacct	300
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<211> 348	
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gtctcacgtt ttagtggaaa tagtggaaagc acattctacg cagactccgt gaaggggccgg	180
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gccgaagaca cggccgttta ttactgtgcg aaagatctgt cgagtgggtc atactactac	300
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gagttggatag ggcgtatcta cggcagaggc actaccaatt acaaccgtgt tttccggaggt	180
cgagtcaatgt ttttgtccagg agtcaatgtt tcttggaaatt gagatgtg	240
accggccgtc acacggccgt ctattactgt gcgagagaca aggggtccga atactctac	300
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<210> 122	

<211> 376  
<212> DNA  
<213> Homo sapiens

<400> 122  
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cattcagcag atatgttatac agctgggtgc gacaggcccc tggacaagggttttgggtccag 120  
tgggagggtat catccctccc taaactacgc acagaagtgc cagggcagag 180  
tcacgattac cgcggacat tccacgaaca cagcctacat gggcttgagc agcctgagat 240  
ctggggacac ggccgtgtat tactgcgcga gagtggccta tgatggtagt ggctattaca 300  
acaatatccc aaagatctac tactactcct acatggacgt ctggggcaaa gggaccacgg 360  
tcaccgtgtc ctcagc 376

<210> 123  
<211> 5  
<212> PRT  
<213> Artificial Sequence

<220>  
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<400> 123  
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